

In the Claims

19. (Amended) A modified kringle 5 peptide selected from the group consisting of NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Lys-NH₂ (SEQ ID NO: 17); NAc-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂ (SEQ ID NO: 18); Nac-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂ (SEQ ID NO: 19); NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Ala-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂ (SEQ ID NO: 20); NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Lys-NH₂ (SEQ ID NO: 21); NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(N ϵ -MPA)-NH₂ (SEQ ID NO: 22); (MPA-AEEA)-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 23) and (MPA)-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 24).

20. (Amended) A modified kringle 5 peptide selected from the group consisting of: NAc-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(N ϵ -MPA)-NH₂ (SEQ ID NO: 25); (MPA-AEEA)-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 26); (MPA)-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 27); NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Ala-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(N ϵ -MPA)-NH₂ (SEQ ID NO: 28); (MPA-AEEA)-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Ala-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 29); and (MPA)-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Ala-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 30).

21. (Amended) A modified kringle 5 peptide selected from the group consisting of NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Lys-(N ϵ -MPA)-NH₂ (SEQ ID NO: 31); (MPA-AEEA)-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-NH₂ (SEQ ID NO: 32); (MPA)-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-NH₂ (SEQ ID NO: 33); NAc-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(N ϵ -MPA)-NH₂ (SEQ ID NO: 34); (MPA-AEEA)-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 35); (MPA)-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂ (SEQ ID NO: 36); NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Lys-(N ϵ -MPA)-NH₂ (SEQ ID NO: 37); (MPA-AEEA)-Pro-Arg-Lys-Leu-Tyr-Asp-NH₂ (SEQ ID NO: 38);